

Sequence listings

SEQ ID NO: 1

Homo sapiens transforming growth factor, beta receptor II (70/80kDa),
complete antisense sequence

Derived from AY675319

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1  CAGCCACACT GTCTTTAACT CTCAGCCCAC CCACACTGAG GAGGGTGCCT AGAGGTTCTA
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241 ACAAGTTTCC TATGGAGTTT TAAAGCTCAT ACTTTGTTCT TGTCTTGTG GACACAAATT
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361 AAGGTTGAGA AGGCTGGCAT GATGGAGAGA ACGGTAACCA TGAGGAAAGC TTCTTGAGT
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481 GGGCTAACTG AGACCCTTAA AGGAGTTCCC CTTTAGTCCA ATAAAAGGCC AACTTCAAAT
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28381	ATGACTTGAG	ACAGGAAGTT	CTTCCTTCCC	CTCAGCTATC	TCTCTTTAGC	ACAGTTTCCT
28441	TCTTTGGATC	TCAGCCAGTT	GCATATATAC	TCTCGCTTTA	GGTGAACATG	CTGGCATGTG
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33961	TGTGTCCATA	CTAGATTCCA	CTCTGCTGAT	TCATACCAGA	ATGAGGTTCT	AGAGTACAAA
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34201	CCAGACATAG	GGCCATATTG	ATGAATTCCA	AGTGTGCTTT	CTAACCAAGG	CTGGTTCACA
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71221	CACCAATAAA	GTACACAGAA	ACCCATAGTG	AACTGAGGTC	TAAGTGGCTG	CTATCAACCA
71281	ACACAATCTT	GAAGTGTCTG	GTCTGATCAT	TTCATTTTCT	TAAAACTTCA	GGACAGTTCA
71341	ATTTAGAGAG	CACAAGAGTG	AAGTTAGCCT	CCCTGGTCAT	CTAAAGCAAC	ATGGGAAGGT
71401	AATTTAAGAA	AAGTGACACC	ATATCAATCA	CTTTCCCAAG	CTTTTCAGGG	GCTCACTCCA
71461	AAGACTTTCT	TGATGATAAA	TTTTATTAAA	GTGCTGATCA	TAATCAAAAC	AGTACATTGT
71521	ATCAGAGGTC	AAGCATTGTT	TTTCACTTCC	CTTCCCAGAA	CTTATTAAAG	ATGTACCTCT
71581	GAAGAAAGAC	ATAAGGAAGA	ACAAAATTGT	TTTGACTCAC	AGAAAATGGC	TTACACAGAC
71641	ATCTAATACT	TAGTCAAGTT	GCTTATATAT	ATGAAATAAG	TTATACCTTG	GGAGTAACAA
71701	AGCCAATAGG	CTTGGCTCAT	AGGGGGCTAA	CTACGGCTGC	AGCAGTTATT	CAAGAAGTTA
71761	ACATATTCTC	TCCTGTCCCC	TGAAGACACC	ACAGGCATCT	CAGTAATAAA	CTGATAACAA
71821	CCAATACTCA	AGAGTTTGTG	AAAATGCTTT	GTAAAGTGCT	GAATAATGTA	AGTGATTATC
71881	TACTTACCAC	AAACGTCCTC	ACCAACATCA	TCATTAAAAT	CTTGAAAGAT	AAGTGCTTTT
71941	CAGTAGTTTT	GAGAGTGAAC	CAATGACATG	CTAAATTAAA	AGCATTATAT	TTCAACCATT
72001	TCCAATAAGT	GCCACAGAAT	ATTAAGACTT	AAGAATATTA	AAATTCTGGT	CAGGTGCAGT
72061	GGCCCATGCC	TGTAATCCCA	GCACTTTGGG	AAGCCGAGGA	GGGTGGATTG	CCTGAGGTCA
72121	GAAGTTCAAG	AGCAGCATGG	CCAACATGGC	AAAACCCTGT	CTCTACTAAA	AATACAAAAA
72181	TTAGCCAGGT	GTGGTAGTGG	GCACCTACAA	TCCCAGCTAC	TCGGGAGGCT	GAGGCAGGAG
72241	AATAACTTGA	ACCCAGGAGG	CAGAGGTTGC	AGTGAGCTGA	GGTCGTGCCA	TTGTACTCCA
72301	GCCTGGGTGA	GAAGAACAAA	ACTCTGTCTC	AAAAAAAAAA	AAAAAAAAAG	AATATTAAAA
72361	TTCTTATTTT	ATCTCCTTTT	CCTCTATAGA	TAGAGCAGGT	AGGGCACTAC	CAGTCCAAAT
72421	CTATGTAATG	TGATGTCAGT	GGTAGCTCCA	GAATTCCCAT	AGAGGAGTGG	CTACATGGGT
72481	GATGATCTGG	CTGAAAGGGA	GGCAGGCAAC	ACAGCTGGAA	GCTGTGCTTG	CACAAGAGAA
72541	CACTACTTGT	GGTACCTAGA	CTGCATGTTT	ATTCAGGATG	GATGGGGACT	ATGGTGGACG
72601	AGAAGGCTGA	AGGCCAAATT	CAAGACCATC	CTTAGAAACC	CACTCCTATT	AAGCACCCAG
72661	GTAAGAAAGG	CCTCATATTT	AAGTCTCCTT	AGCATGTATC	ATTTTAACAG	GGCTTTCCAA
72721	AAAGTACAAC	CAGGGGTCAC	CTTTGCAATA	TGGGATTATT	TACCCTCAGG	TAGGTAGAAG
72781	AGTATCTTCA	GCCACTCTTC	ATTGCACCAA	GACCCCTTCA	GCCATCCCTA	TTCCAGCAAT
72841	CCCAATGACT	GAAATCATCT	ATGACTCATA	AAGAGAAAAG	GCAGGAGTGA	CAAAAAATTG
72901	TCCTGCGTGA	GATGGTGTGA	TTTACTTTAT	GACTTTGCCT	GGTCTCTATA	GTCATAATGC
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73141	TACGAGATTT	TCCATAAGCC	TATGGTTCTG	TCAAACAATA	CTACAGGTGG	CTTTAAGCCA
73201	TTGACATTTA	CACTTGGTGG	TGAAAAGCCA	TCTGGGAGAA	AAAAAAATAG	CACTCCCCGC
73261	CTTGGATCTA	AAAGCAAGTT	TCCTCGGCCT	TTAAAAGCAA	TGAATTCCAA	ACAGCAACAA
73321	ACAAAACACA	TAAACAGAAT	TAGTGAGAAT	CTGTGCTAAG	AGCTATGGCA	AAATCAGACA
73381	GAAGTTTCTC	CCTCAAAGAG	TTTTCATTTT	AGTAAGAGTT	TTGTTTTGTT	TTGTTTTGTT

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73561 CTACAGGCAC CCGCCACCAC GCCCAGCTAA TTTTTTGTAC TTTTAGTAGA GACAGGGTTT
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76321	AAAGTTGAAA	GATTCCTAGT	CTACGCTAAC	AAGTGTCTGC	AAACTCTACA	GAAATGCAAT
76381	TAGAGGTTGC	GGCAGCTACT	CCCCTGCCTA	AAACAGCAGT	CTGAAAACCTG	CCAATCTGTT
76441	GCAAATTCTG	TCTTTTCTGA	GAATATTTTA	AGAAAAGTGG	TAGAGAAATA	TTTGAAAGGC
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77101	ACAGCCTGGT	CACCCAGACA	CCTAGATTCC	TGGTACAACA	TACCTTAGTC	TTTACCAGCT
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77581	ATACAGGCCG	GGTGCAGTGG	CTTACACCTG	TAATCCTAAC	ACTTTGGGAG	GCAGAGGTGG
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Homo sapiens transforming growth factor, beta receptor II (70/80kDa),
antisense sequence

Derived from BC040499

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LOCUS AY497473

DEFINITION Homo sapiens transforming growth factor, beta receptor I
(activin A receptor type II-like kinase, 53kDa) (TGFBRI) gene

coding sequence in reverse complement

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1081 CATCAACATG AGTGAGATGC AGACGAAGCA CACTGGTCCA GCAATGACAG CTGCCAGTTC
1141 CACAGGACCA AGGCCAGGTG ATGACTTTAC AGTAGTTGGA AGTTCTATTT TATTGCAATG
1201 GTCCTGATTG CAGCAATATG TTGTAGTCAC AGACCCAGTT TTTGAAGAGG GTGCACATAC
1261 AAACGGCCTA TCTCGAGGAA TTAAGTCAAT TTCAGCTATA CACATGCTGT TGTGTATAAC
1321 TTTGTCTGTG GTCTCTGTGA CAGAGACAAA GCAGAGCCCA TCTGTCACAC AAGTAAAATT
1381 GTCTTTTGTA CAGAGGTGGC AGAAACACTG TAACGCCGTC GCCCCGGGA GCAGCGCCGC
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1441 CGCCGCCGCC GCCGCCGCCG CCAGCACGAG GAGGAGCAGC CGGGGACGCG GAGCAGCGAC
1501 CGCCGCCTCC AT

SEQ ID NO: 95

LOCUS AY497473

DEFINITION Homo sapiens transforming growth factor, beta receptor I
(activin A receptor type II-like kinase, 53kDa) (TGFBRI) gene

mRNA in reverse complement

1 AAACACTATA GGTTTTAACA TAGAGACTTT CAGCAATTCT ACTCATTTCC ATTAGAAAGA
61 CACAGAAGTG GCACTTACTG GTATAGTACA ATCCCATTTT GAAGGCATGT AATGTTCTGA
121 ATAAGTGAAA GAACAATAAT AGTATACAAA ATACAATTGC ATGAATTATG TTCCTCACTA
181 CTATATGAGG TCATTTTTAG ACTCAAGATA AGAGTTTTAT AAGTGTTAGT TTTAAGATCC
241 TGAAAAACTA TAGAACAGTA CATTCAAAGT CTGAATCAAG GAAACTCTAG TGGTTCAGAA
301 TCCTCTGAGA ATGTACTAAC CAGGAGTAAA TCACTTCTT TAGTAATAAG ACATGTTTCA
361 TTGTAATTCA GCAATCCAAC TCCTTTGCCC TTAAAGATGA TCTCCAGCAC AGCAGAGTTA
421 CCTAAAGTTA AGACCAGCAA TCATCTTTTT AAAAACAAG TTTTGTTAAT AAAAAATAAA
481 GGTAGACTAC ACATTTTCTG TCCTGGGAAA GAAGCGTTCA TAGTGCACAG AAAGGACCCA
541 CATGGCTGTT TCCTGGGTCC AAAGAAATCC TGGGAAGTTT TTAATTGACT TTATTACACT
601 GCTGCAAAAG GAAGCAATAT CCTTCTGTTT CCTCTCAGTG AGGTAGAACA ATTGACCTCC
661 CAAATTAAAA CCCAGGAGCA GATCTGAAGA AAAAAGGAGA GTTCAGGCAA AGCTGTAGAA
721 TTACATTTTG ATGCCTTCCT GTTGACTGAG TTGCGATAAT GTTTTCTTAA TCCGCAATGC
781 TGTAAGCCTA GCTGCTCCAT TGGCATACCA ACATTCTCTC ATAATTTTAG CCATTACTCT
841 CAAGGCTTCA CAGCTCTGCC ATCTGTTTGG GATATTGGC CTTAACTTCT GTTCACAAAC
901 AACTTTTCTC ATTTCTTCAA CTGATGGGTC AGAAGGTACA AGATCATAAT AAGGCAGTTG
961 GTAATCTTCA TGAATTCCAC CAATGGAACA TCGTCGAGCA ATTTCCCAGA ATACTAAGCC
1021 CATTGCATAG ATGTCAGCAC GTTTGAAGGA TTCAAATGT TTCATATTTA TGGAATCATC
1081 GAGAACTTCA GGGGCCATGT ACCTTTTTGT TCCCACTCTG TGGTTTGGAG CAATATCAAT
1141 GGTATCTGTG GCTGAATCAT GTCTTACTGC CAGTCCTAAG TCTGCAATAC AGCAAGTTCC
1201 ATTCTTCTTT ACCAAGATAT TCTTTGATTT CAAATCTCTA TGAGCAATGG CTGGCTTTCC
1261 TTGGGTACCA ACAATCTCCA TGTGAAGATG GGCAAGACCG CTCGCCGTGG ACAGAGCAAG
1321 TTTTATCATT CCTTCCACAG TAACTGTGTA TCTGTTTAAG TAATCAAAAA GGGATCCATG
1381 CTCATGATAA TCTGACACCA ACCAGAGCTG AGTCCAAGTA CCATTGTCTT TATTGTCTGC
1441 TGCTATAAAT CCCAGGATGT TTTCATGACG TAACATTACA GTTTGATAAA TCTCTGCCTC
1501 ACGGAACCAC GAACGTTCTT CTCTAGAGGA GAATATCTTA ACAGCAACTT CTTCTCCCCG
1561 CCACTTTCCT CTCCAAACTT CTCCAAATCG ACCTTTGCCA ATGCTTTCTT GTAACACAAT
1621 AGTTCTCGCA ATTGTTCTCT GAACAAGCAA TGGTAAACCT GAGCCAGAAC CTGACGTTGT
1681 CATATCATAA ATTAAGTCTT TCAACGTAGT ACCCTCTGAA ATAAAAGGGC GATCTAATGA
1741 AGGGTCCTCT TCATTTGGCA CTCGATGGTG AATGACAGTG CGGTTGTGGC AGATATAGAC
1801 CATCAACATG AGTGAGATGC AGACGAAGCA CACTGGTCCA GCAATGACAG CTGCCAGTTC
1861 CACAGGACCA AGGCCAGGTG ATGACTTTAC AGTAGTTGGA AGTTCTATTT TATTGCAATG
1921 GTCCTGATTG CAGCAATATG TTGTAGTCAC AGACCCAGTT TTTGAAGAGG GTGCACATAC

1981 AAACGGCCTA TCTCGAGGAA TTAAGTCAAT TTCAGCTATA CACATGCTGT TGTGTATAAC
2041 TTTGTCTGTG GTCTCTGTGA CAGAGACAAA GCAGAGCCCA TCTGTCACAC AAGTAAAATT
2101 GTCTTTTGTA CAGAGGTGGC AGAAACACTG TAACGCCGTC GCCCCCGGGA GCAGCGCCGC
2161 CGCCGCCGCC GCCGCCGCCG CCAGCACGAG GAGGAGCAGC CGGGGACGCG GAGCAGCGAC
2221 CGCCGCCTCC ATGGTCCCGC CGCCACCGCC TGTGGCCCGG CCCGGCCCGG CCGCGCCGCT
2281 GCCTCACCCC AGCAAACCTC GCCTCGCC

SEQ ID NO: 96

LOCUS AY497473

DEFINITION Homo sapiens transforming growth factor, beta receptor I
(activin A receptor type II-like kinase, 53kDa) (TGFBRI) gene

gene in reverse complement

1 CCTAGGATGG GCAGTATAAA CAACTGTGTT TTAAATGAC AAAATTATTA CTCTAATCAA
61 ATTTATCTGC AGTTATCGAC AGTAATCATG ATCCATCCCA CCTTTTAAGT TTCTTTCATT
121 TTCAATATCA GCCAATTGTG GTTTATCAAA CCATCCCTAG CCAAAAAGG AATTAAGTGC
181 AGTAATACGT AAAGTCAATT TAGAACTAAA CTCCAACCTT AAAAAAATTC CCTGATTCAT
241 CAAAAGTACA TTTTATCCAC TTAGGTAAGA AAAGTATAAT TGTAAGTCAA TCTTTGATTC
301 AATAAAAAAT ATCTTCAGCA AAAGTGGATA ACCTTACATG AAAGTGAGAA ATCATGTATT
361 ACAACTTTAA ATCAGACAAC TTTTGGTATA TTCTTAAAT TATATAAAAA TAGTTTTAGT
421 GATAAAATGC ATTGGTCCCC TGGATTTTCAT AAAGCAGATC TGGTAGGCTT AGAAATGGCC
481 CAAAACAAC AACAAAAAAA AAAAAACAAA AAAAAACCC CCACAAAAAA AGTACCTGAT
541 CACAGAATTA GAGATTATTA TTAAACAAGT AATACAACAC AACTATAAAA ACAACCTCT
601 CAACTCATT CATTTGCTTC AATGAAATTA TGGTAACTAC AGAAAAAAC TCAACTTTGT
661 AGTCAAATA GTAACCTCTA AGCATTTTTC CCATGATGTT TTAAACACA TGAAATTGAC
721 ACTAAATTCC TATGTAATAT GTCACCTATT CTAAAATTAC TTTGGTTTTA AGATCAAATA
781 GTCTTTCAAA TTTATTCTAC GTATAAGAGA TAAGACATTA TCAGAATGCA ACTCATTCAG
841 TGTAATGGC CACTTCTCTC TCCTGACCAA AATACTGTAT GGGTTACTCT GACCCTGTAT
901 CGGGGAAGT ATAGCACTAT AAAGTCGAAG TCTTGCCCTCA GAGCTATGAT AGTTTTTCT
961 CCCCTCAGAA TAAGATCACA GTGATAAAAG GACTTCGAAA ACTGTAAAAG CCCTGCAGAG
1021 ACTTCATAGC ATTGGAAGTA AAAACCAAGT ATTATAAGCT AGTTTCAACT TAATGTAAGA
1081 AGACCATGAC AAGTTTGCTT TCAATATTTG ACCAAAGACA TCTGTAAATG GAGAAAAACC
1141 TATACTCAGA ATGTTCTTTA GCTACCACCT CTCCCAAGTA ACAAGGGTGC ACTGAATGCA
1201 TTACATAAAT TACATGTAAG CAGTTGCAAT GTACGCACAC ATAAGGAAAC ACTGAATTAA
1261 AAGCTGCCTT CCTCAAAGTG TAGTGAGACC TCAAAACCAC ACTAGCTCTA ACCTCAGTTA
1321 CAAAGACAGA GGATCCACCG AACGTTTACT CCTACATACA TAACAAATGT GACTTTCTTG
1381 CACATCTGTT ATCTATTCAA ATTATTAAGT ACCTTCGCCT TCCTAGAAAA AGGCCTATTA
1441 TTTAATATTA ACCATATTCT AATTTTAAAT GTAATTCATG TCTTAAAAAA TAAGTCTTTG
1501 AACAGATAAA GAAAAAATG GCTTACAGAA TACCACATAA CTTTTCACAA GCAGCTAGAC
1561 AGACTTTCCC TTTTAGAAAA ATTAGTCTGT ATGCTACAAA TATTAAAGTA CTATAAGCAT
1621 CTAGAGTGCC ACTTGATTTT TAAAAGTAT ACTTAGACAA CACAATTAAA ATGCAAAAGC
1681 TTGATGTGAG AATATTCAAA CATGACCATG CTAATAATTT ACTAATTAAT GCCACTTCAT

1741 TTCTTTAGTG GCTTAAGCAC ATTTTGGTAG GAGAAAGGCA TTTTCAGAAT AGAATACCAA
1801 ATGACATACC ACAAAAACCC TTCCAAAAAC AAAACAGAAA AAGTTTGGGT TACCCAGATA
1861 AATAAGCTTT TTGAATTCAA ACACTATAGG TTTTAACATA GAGACTTTCA GCAATTCTAC
1921 TCATTTCCAT TAGAAAGACA CAGAAGTGGC ACTTACTGGT ATAGTACAAT CCCATTTTGA
1981 AGGCATGTAA TGTTCCTGAAT AAGTGAAAGA ACAATAATAG TATACAAAAT ACAATTGCAT
2041 GAATTATGTT CCTCACTACT ATATGAGGTC ATTTTTAGAC TCAAGATAAG AGTTTATATA
2101 GTGTTAGTTT TAAGATCCTG AAAA ACTATA GAACAGTACA TTCAAAGTCT GAATCAAGGA
2161 AACTCTAGTG GTTCAGAATC CTCCTGAGAAT GTACTAACCA GGAGTAAATC ACTTTCTTTA
2221 GTAATAAGAC ATGTTTCATT GTAATTCAGC AATCCAACCT CTTTGCCCTT AAAGATGATC
2281 TCCAGCACAG CAGAGTTACC TAAAGTTAAG ACCAGCAATC ATCTTTTAA AAAACAAGTT
2341 TTGTTAATAA AAAATAAAGG TAGACTACAC ATTTTCTGTC CTGGGAAAGA AGCGTTCATA
2401 GTGCACAGAA AGGACCCACA TGGCTGTTTC CTGGGTCCAA AGAAATCCTG GGAAGTTTTT
2461 AATTGACTTT ATTACACTGC TGCAAAAGGA AGCAATATCC TTCTGTTCCC TCTCAGTGAG
2521 GTAGAACAAT TGACCTCCCA AATTAAAACC CAGGAGCAGA TCTGAAGAAA AAAGGAGAGT
2581 TCAGGCAAAG CTGTAGAATT ACATTTTGAT GCCTTCCTGT TGAAGTGT GCGATAATGT
2641 TTTCTTAATC CGCAATGCTG TAAGCCTAGC TGCTCCATTG GCATACCAAC ATTCTCTCAT
2701 AATTTTAGCC ATTACTCTCA AGGCCTACAA GAAAATATAA AAAAAAAAAAT TAATGCATGC
2761 ACCATTTTCC ATTGGTCTGG ATAAGAATTT TTACGTGTAA ATTTTCTTTT TAAATGACAT
2821 ATACACTTGT TATTGTTTAA GCATTATTTG TAATAGCAAA TGAGTAGAAA TTACATAAAT
2881 TTCCATCAAT AGGGTACTGG TGCCAGTGCT GGAAAGCAGG TGGACAAAGA GTGACTCATT
2941 TAGCAAACCT GAAAAGGCCG AATCCTAAAC CAGAGCTGGG CAAAGCAAAG AAGAAATCTG
3001 ACCTCACACT GAAATTTTAA GTTTGTACTT TGAAGTATTC ACATCAAATT TTTCTTTCTT
3061 TTTTTTTTTT TTGAGACAGG GTCTCACTCT GCCGCCCAA GTGGAATGCA ATGGCATGCC
3121 CTCAGCTCAC TGCAGCCTTG ACCTCCCAGG CTCCAGCAAT CCTCCCCTT CAGCCTCCCA
3181 AGTAGATCAA ATTTCTAAAA GACAAAATCT AGCAGCACTC TCATTTATTA AACTATGTCA
3241 CATGCAAATT CCTTAAGCAT GCTGAAATAC ATTTTATTTA TTTGGGCCTG GAGATGTGAT
3301 TTTGTTGGAG GTAGCAAGCT CTCTTGCTTA TCAGAGGTTC TGCTGAGTCC TACAATAGTT
3361 TAAGACGTAA AATCAACTAC TGAAGTGAAC AGCTTTTTAT AGCCTTTTTA AGCTACAAAT
3421 CTTCAAACCT GGAGGAATAT GCTCTGCTTC ACCAATATTG TAATAGTAAC AGCTAGCAAT
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3541 TTCTAAACT TTTTGTTAGA GCTGAAGTTG AGCAGCTCAG ATACTGCTGG AAAAAAGCAT
3601 TTTCCCTAAA TTCTGGTTA ACGGTTATCT GGGCAATCAG ATGAATAAAA CGGAAAATCA
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3961 AATATGAAAA CCAAATTATT CTGTTTTTAA ACATTGGTTT GACTGCTATG AAAAAGAAAA
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4141 TGGGATATTT GGCCTTAACT TCTGTTTACA ACAAACCTTT CTCATTTCTT CAACTGATGG
4201 GTCAGAAGGT ACAAGATCAT AATAAGGCAG TTGGTAATCT TCATGAATTC CTGTATCAGT
4261 TTAAAAAAA TTAAATTTTG ATGAATTTAT TCAGCTAATG CCAAGGCATT AAAAGATCAC
4321 CTGAGAGACC TATTACAGTG TTTCACTACT AAAGTAGGTA TGTGGAACAG AGATTACAGC

4381 TTTCTCTATC TCATCCACAT CTGTATCCAC AAGCCACTTC CCTGAGTTGA TATTCCACCC
4441 ACACCTTGAA ACTAAGTATT GCCATGAGAT CAATAAAGCC TATTTGAAAT GACAGTAAAC
4501 TATGAGCTGT GATGACCAGA TTGACATTTT CCCCTAGAAC CCTATTGTGC AGCAAATGAG
4561 ATTCTATTCT GTAAGTGCAC AGGCTCTCAC TTGCACAGGG ATATATATAC CCTATCAGTG
4621 AGACTTGAAT CTTATGAGGC AAGTTTAAGG TGCCAAGTCC CTTTCTCTAC TTGCATCTGT
4681 CAATAATATA TCCTATTTAC TACAATACAG AGAAATACTC TGTCCCTCAC TGCTTCAAAA
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4921 AAATAACCAC AATCTCAAAG AAAAAAATA AGTTTGACAC CCAGGTACAT GCTAGAAACA
4981 GCTTCCTAGG TGAGCCTCTC AAGCAGCTGT CCATTCTAAA GAACTATATT CCAATATAAC
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5101 GCTAGAGCTG TGCATGTGCA CACCATGGCA CTGTCCCCTC CCAAAGGTTT CATCGCCTTT
5161 GTTTTCTCTG GCACTCGGTG ACATCCTGTT TCAGATAATG GACTGTACAT AATTTCATTC
5221 AATTAAAAAA AGAAGACAAT TCTTGAACAA CTTCTGCTCA TGACAAACTA CTGGGGGAGA
5281 GGAGAGCAAT TTACCACCAA TGGAACATCG TCGAGCAATT TCCCAGAATA CTAAGCCCAT
5341 TGCATAGATG TCAGCACGTT TGAAGGATTC AAAATGTTTC ATATTTATGG AATCATCGAG
5401 AACTTCAGGG GCCATGTACC TAAAAAAAAT TTGCAAAAAG AACTTTGAAA ATCATCCCCT
5461 TACTGCCATA TTTGGATGAA CCTCCTTTCA GACATTTCTT TATACATAGA CATATGTACA
5521 CAATATTACA TAACAAATAT TAGACTCAAG AAAAATATTA AATATCCACA CTGGTTTGCA
5581 CATTAATGAT GTCAACTAAA TCCCACAAAA TGCTAAAATA GCAGTTTCAC ACTATCTGCC
5641 TCATGATCTT GGTTTCCAAT GAAAGACATA CATAACAATT GTGACTGTTA AAGGTGTCCT
5701 AATGTACACC ACCTATACAC GTACTAGGTA GAAACCATGC ATTTTAATTT AGAAGTCTTC
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6121 AGACAGTAGT TTGAAAAAAT GCTGTGACAG CTTGTGATAA GGGGTTTGGG TCAGAAATGC
6181 CCAGGTTCAA ATCTCAGCTC TTCGGCTTAT CATCCACATG AAGCTGGCCA TATTATTTAA
6241 CCTCCTCAAG GTGGTTTCCT CATCTGTGAA TTAAAGGGTA ATAACACCTA CCTAATTAAG
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6361 CTGTGATGGA TGCTGGGAAT ACAAGGCTGC TCTCAAGCAG CTTGCACCTA ATAGAATATA
6421 TAGGCAAATA AATAAACAGT TACAACATAA CGCATAAGTG CCTCAACGGT CCCAGTTCAG
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7141 TTTCAAATCT CTATGAGCAA TGGCTGGCTT TCCTAAAGAA TCAAAAATTA AACATGACTG
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7441 ATCTGTTGGT GCCTTATGAA TTCTTTTAA GAACTTAGC CATAGACTGA TATTCCAGCA
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7621 TGGAAAACCT GGAACCTTCA TACACTGCTG GCAGGAATGT AAAATGGTTC AGGCACTTTA
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7981 AGCCAGTCAC AAAAGACCAC ATATTGTATG ATTCATTTAT ATAAAATGTG CAGAATAAGC
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8101 GAGTGACTGC TAGTGGCTAC AGGGTTTCTC TTTGGAGTGA AGATATTCTA AAATTGACTG
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8221 AGTGGGTGAA CTTTATGATG TGAATTATAA CTCAATAAAA CTGTTACAAT CAATAATAAT
8281 AATAACTACA AAGATGGAGC TAAGATGTGA ACTAGCACTT ATGCTGTATG TAAAGTGGGG
8341 AGTGGGGGAG CCATTGGGTT GCAAGTGGAC AACTGCTAGA GCAGAGAGAT ATACTGGCAT
8401 CTGAAATCCC ACATGCACAA AACTACAGC ATACAAATGT CATAACTAAA ACTTAGTATT
8461 TAATTTTTCA ACTGTAAAC TATCTCATGC AGTCTCCTTA TCTCAAGGTC AAATATAATT
8521 ATTAGATGGC CAGATAACAG TTACCTAGCA TGTTGATGTT TACAAAGCAC TTTCACATTT
8581 CATTTGATAC TACACATAAA AATATATTCA GTTGGACCTG AATTCTGAAA CTGAGATCCT
8641 CAAAATCAT AAGATTGAAT GCTACTTTTA AGAACTCAA AAAAAATCACA GTTTTGTTAC
8701 AACGACTAGA GAATCTCTAA ACAAATTTCC AGATTCCTAG CTCTACCTGA AATTTTGAAC
8761 TGAAAGATAA AATGTTCTCA TCTAAAAAAT TTGTTTTAAA TATCTCCAGT TAGGCCACCT
8821 ACTGTTTTTG TTTTGTGTTT TTTTCTGAG ACAGGGTCTC TCTCTATTAC TCAGCTGGGA
8881 GTGCAGTGGC ATGATGTCTG CTCCTGCAG CTTGACCTC CCAGGCTCAG GTAATCCTCC
8941 TACCTCAGCC CCCTGAGTAG CTGGGACTAC AGGTGCATGC CACCATGCCC GACTAATTTT
9001 TTGTATGTTG GTAGAGATGC GGTTTTGTCA TGTTGCCAG GCTGGCCATG AACTCCTGAG
9061 ATCGGGCGAT CTGCCCACCT CAGCCTCCA AAGTGCTGGG ATTATAGGCA TGAGCCACCA
9121 GGCCAGCCT CCACCTTCTA TTTTCATAGA CATTATTCAT GAAAATTTAA AGCTTAAATA
9181 ATAGAACTGC TTATAGAATT ACCTTGGGTA CCAACAATCT CCATGTGAAG ATGGGCAAGA
9241 CCGCTCGCCG TGGACAGAGC AAGTTTTATC ATTCCTTCCA CAGTAACTGT GTATCTGTTT
9301 AAGTAATCAA AAAGGGATCC ATGCTCATGA TAATCTGACA CCAACCAGAG CTGAGTCCAA
9361 GTACCATTGT CTGTAAAACA GAATTAACAT TTCGGTTGGG CTGCAGACCA TTTATGATTT
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9481 CCACCTTAAG TCTGTATTTT CTTACATTT TCTTCTACAT TGTTTGTTTT ATACTTTTAT
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9601 AATCCCATTT AAGAGACAAA GATGGTGACT TTTTGAGTC TTCACAGATA ATTGGTATCA

9661 GAGCTGGCAC TCTTAAGTGC CTAGAAAAAT CTCTGGGTCG GGTGTTGGTGC TCAGTGGCTG
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9781 GAAGTCTCCG GAGACCAGTG TGGCCACAGT GGGAATGGAA GAGTATGGGA CAAAAGTGA
9841 AGTGAGTTTC AACAAGAAAT GACAAGCTTT AGTGACAGAT TGTGTACACA GTTCACAAAG
9901 ACGCAAGACA AGATGATTAT GATTCCAAGG TTGCTAGCTT AGAAAAACAG AAATAGTGAT
9961 ACCTCTAACA CATGGGGAAG CTGAAATAGG AACTGAAATG TCAGGAGCAA AACTGGAAAA
10021 GCTGGGACTA AACTCAAACA ACGAGGAACG TGGGTTCCAA AATTTCTCTT GTAAGCCAAA
10081 ACACGAGTTT AAAACATGTT TTCTCATAGG AATAATGGTA AAGACACTAG CTAGGCTCCT
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10201 TAATTAAGCC ATGAAGAAAA AAAGCATTAC AACATTAGGA GACAGCACAA TCCAGTAAAA
10261 GATGGACGCT TAGAGTCATT CTAACCTGGT TGGAAATCCAC GGCCTGCCTG TTAGTGACAA
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10501 GTATTTGGTT GAAGAAGAAA ACCAACTCTG ATTAGAAGAG ATTAAGAAAG GAATAGAAAC
10561 TTTTCAGAAG ATTCTGGAGG AGATTTCTGG GTACATTTAA AAGGACTTAG TATTAATTTG
10621 TTCTTCATGA ACTTTATAAA CTGATTCTGA CCAAACCTCGC ACCCTTGCTT TTCCTTGATA
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10741 CAAATACAAG AAACAAAGTG GAAGACAGGC AGATGGAAGA AAGGGAAACC ACAGGTTATG
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10861 GAATTAAC TAATCAGGAGA AGATAAAGGA ACTGAGGTGG AGGGGCTAGG CTCAGAGAAA
10921 GTACATTAAC CCCTGACAAA GAAGAAAGAA GGGGAGTAGG GAAGAGCCAG TAGGGTCAGC
10981 TAGTTCCCAT GGTGACCACC AGATGACAGC ATCAGCCTGG TTAGAAGTTC AGGGCACAGA
11041 CTCACCAACA TTTTGTGCTG CTGTGTTTGT AACTTGGAAT GTGACAAAAT CCAGACCCTG
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11161 CATGAATGTG GCTATGTTCT TAAGGAAGAA ATTACCACTT ATAAATCAG TACCTTCATA
11221 TGAACCAGTG AGAAGAGTAA GAGTTGAGAA AACAAAGACA ACTATGAAGA AGTAAGAAGG
11281 AACAAGGAAA GTCTAGGGTC ATCAAACACA GTAAATAAAG AAAAGGTTGT TTTAAGAGTT
11341 GGGGGAGGAA ATAAATGGTG ATAAAGGCAG AAACGAAAAT GAGAACTAAG GATTTGGTCA
11401 GACTGATTAC TATTTTTTGT TTTGTTTTGT TTTGTTGTTG TTTTTTTTTT GAGACAGAGT
11461 CTCCCTGTGT CGCCCAGGCT GGAGTGCAGT GGTGCGATCT CAGCTCTGAC TGCAACCTCC
11521 ACCTCCCAGG TTCAAGCAAT TCTCTGCCTT AGCTTCCCAA GTAGCTGGGA TTATAGGCAC
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